

1 **Effects of Surotomycin on *Clostridium difficile* Viability and Toxin Production In**

2 ***Vitro***

3

4 Laurent Bouillaut<sup>1</sup>, Shonna McBride<sup>2,3</sup>, Joseph A. Sorg<sup>4</sup>, Diane J. Schmidt<sup>5</sup>, José M.

5 Suarez<sup>2#</sup>, Saul Tzipori<sup>5</sup>, Carmela Mascio<sup>6</sup>, Laurent Chesnel<sup>6</sup> and A. L. Sonenshein<sup>1\*</sup>

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7 <sup>1</sup> Department of Molecular Biology and Microbiology, Tufts University School of Medicine,

8 Boston, MA USA; <sup>2</sup> Department of Microbiology and Immunology, and <sup>3</sup> Emory Antibiotic

9 Resistance Center, Emory University School of Medicine, Atlanta, GA USA; <sup>4</sup>

10 Department of Biology, Texas A&M University, College Station, TX USA; <sup>5</sup> Department

11 of Infectious Disease and Global Health, Tufts University Cummings School of

12 Veterinary Medicine, North Grafton, MA USA; <sup>6</sup> Cubist Pharmaceuticals, Lexington, MA

13 USA

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15 Running title: Effects of surotomycin on *C. difficile*

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17 \* Corresponding author: A. L. Sonenshein, Department of Molecular Biology and

18 Microbiology, Tufts University School of Medicine, 136 Harrison Avenue, Boston, MA

19 02111. Tel: 617-636-6761; Email: linc.sonenshein@tufts.edu

20

21 # Current address: C/ Núñez de Balboa 54, 28001 Madrid, Spain

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23

24 **ABSTRACT**

25 The increasing incidence and severity of infection by *Clostridium difficile* has stimulated  
26 attempts to develop new antimicrobial therapies. We report here the relative abilities of  
27 two antibiotics (metronidazole and vancomycin) in current use for treating *C. difficile*  
28 infection and a third antimicrobial, surotomycin, to kill *C. difficile* cells at various stages  
29 of development and to inhibit the production of the toxin proteins that are the major  
30 virulence factors. The results indicate that none of the drugs affects the viability of  
31 spores at 8X MIC or 80X MIC and that all of the drugs kill exponential phase cells when  
32 provided at 8X MIC. In contrast, none of the drugs killed stationary phase cells or  
33 inhibited toxin production when provided at 8X MIC and neither vancomycin nor  
34 metronidazole killed stationary phase cells when provided at 80X MIC. Surotomycin, on  
35 the other hand, did kill stationary phase cells when provided at 80X MIC, but did so  
36 without inducing lysis.

37

38 **INTRODUCTION**

39

40 In the United States, the incidence of *Clostridium difficile* infection (CDI) has been  
41 steadily rising over the last 15-20 years (1-3). There has been a significant increase in  
42 primary, recurrent and untreatable CDI on a global scale as well (3, 4). Treatment with  
43 oral vancomycin or metronidazole, the current standard practice, leads to recurrence of  
44 CDI in up to 30% of patients treated for an initial episode. For patients that have already  
45 suffered multiple recurrences, the future recurrence rate can be as high as 60% (3, 5-7).

46 In addition, many cases of CDI in the US are now linked to a family of more virulent  
47 strains (NAP1/BI/027) that emerged first in the UK and then in Canada (8, 9).

48

49 *C. difficile* spores, the infectious form of the organism, are metabolically dormant but  
50 germinate in response to a combination of host-derived bile acids and glycine (10).  
51 Because spores are metabolically dormant, they are insensitive to the action of most  
52 antibiotics. Thus, after antibiotics are prescribed to treat the active infection, the spores  
53 that remain in the colon or that are acquired from the environment are able to germinate  
54 and re-initiate active infection. The process of spore germination is largely enzymatic  
55 and, once initiated, cannot be stopped. Once stimulated to germinate, the spore  
56 releases a large depot of dipicolinic acid, chelated 1:1 with  $\text{Ca}^{++}$  (CaDPA), in exchange  
57 for water (11). Also, the cortex layer, a specialized peptidoglycan, is degraded (12, 13).  
58 After these changes, the spore is no longer dormant and does not have the resistances  
59 typically characteristic of a spore. Metabolism then resumes within the spore core and  
60 a vegetative cell grows out from the germinated spore. Such outgrowth is susceptible to  
61 the action of several antibiotics (14). Moreover, the host intestine contains an  
62 assortment of antimicrobial compounds that are produced by the innate immune system  
63 and the host microbiota. To combat these defenses, *C. difficile* can modify its cell wall to  
64 resist antimicrobials or efflux a variety of antimicrobial peptides from the cell (15-17).  
65 These resistance mechanisms allow *C. difficile* to grow in the presence of antimicrobial  
66 peptides and may also contribute to resistance to therapeutic antimicrobials.

67

68 Given the difficulties in treating *C. difficile* infections, the identification of narrow-  
69 spectrum antibiotics that are potent inhibitors of *C. difficile* growth would fulfill a critical  
70 need. Also needed are compounds or methods of reducing spore abundance or  
71 viability in order to limit the onset of infection and to prevent relapsing disease. Here we  
72 investigated the activities of a newly-developed antibiotic, surotomycin (18, 19), and  
73 compared its activity to that of metronidazole and vancomycin, two antibiotics that are  
74 currently used for treatment of *C. difficile* infections. Surotomycin is a lipopeptide  
75 antibiotic that acts through depolarization of the membrane, leading to the loss of a  
76 proton gradient and cell death (19). The mechanism of action of surotomycin differs  
77 significantly from that of vancomycin (cell wall synthesis-inhibiting) and metronidazole  
78 (inhibition of cellular enzymatic functions), suggesting that surotomycin may affect *C.*  
79 *difficile* physiology differently than do these other antibiotics. We report here the relative  
80 sensitivities of *C. difficile* strains to surotomycin, metronidazole and vancomycin at  
81 multiple stages of the life cycle, i.e., as dormant (phase-bright) spores, as germinated  
82 (phase-dark) spores, as exponential phase cells and as stationary phase cells. In  
83 addition, we assessed the effects of the antibiotics on toxin gene expression and  
84 measured the effects of various mutations that affect the sensitivity of exponential  
85 phase cells to nisin (an antibiotic that creates holes in membranes and disrupts cell wall  
86 synthesis) on susceptibility to the three tested compounds.

87

## 88 MATERIALS AND METHODS

89

90 **Strains used and growth conditions.** *Clostridium difficile* strains UK1 (ribotype 027;  
91 provided by Dr. D. Gerding) (20) and JIR8094 (ribotype 012; provided by Dr. J. Rood)  
92 were routinely grown at 37°C in brain heart infusion (BHI) medium supplemented with  
93 cysteine (0.1%) and yeast extract (0.5%), referred to here as BHIS, or in tryptose-yeast  
94 extract medium (3% tryptose, 2 % yeast extract, 0.1% thioglycollate), referred to as TY.  
95 All growth experiments were performed in Coy anaerobic chambers in an atmosphere  
96 containing 85% nitrogen, 10% hydrogen, and 5% carbon dioxide as previously  
97 described (21, 22). Susceptibility testing of surotomycin requires the presence of 50 mg  
98 (2.5 mEq)  $\text{Ca}^{++}$  per liter in the medium (23). The levels of  $\text{Ca}^{++}$  in the media used were  
99 determined by Laboratory Specialists Inc; following their instructions, the  $\text{Ca}^{++}$   
100 concentration in all media was raised to 50 mg/L by addition of  $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ . Drugs were  
101 added where indicated. Surotomycin was provided by Cubist Pharmaceuticals.  
102 Metronidazole and vancomycin were purchased from Sigma-Aldrich. Nisin was obtained  
103 from MP Biomedicals.

104

105 The following mutant strains derived from JIR8094 were used in this study: MC112, a  
106 *lytC* (peptidoglycan hydrolase), nisin-tolerant mutant; MC119, a spontaneous *cprK*  
107 (nisin-resistant) mutant with increased resistance to lantibiotics and polymyxin B (16);  
108 MC120, a mutant with a TargeTron insertion in *dltD* resulting in increased susceptibility  
109 to nisin and polymyxin B (15); and, MC141, a mutant with a TargeTron insertion in *cprA*  
110 causing increased susceptibility to lantibiotics (16).

111

112 **Minimal Inhibitory Concentrations.** Minimal inhibitory concentrations (MICs) were  
113 determined by growth of bacteria in culture tubes or in 96-well plates (that had been  
114 pre-reduced for at least 24 hrs prior to use) in media (either BHIS or TY) supplemented  
115 with 2-fold serial dilutions of the test compounds. The initial bacterial titer was  $5 \times 10^5$   
116 colony-forming units (CFU) per ml. Each strain and each drug concentration was tested  
117 in duplicate in each MIC assay. The MIC was determined as the lowest concentration of  
118 drug at which no bacterial growth was detected after 18-24 hrs at 37°C. MIC assays  
119 were performed at least 3 times to ensure reproducibility of results.

120

121 **Spore germination assay.** Spores of *C. difficile* strain UK1 were produced and purified  
122 as described previously (14, 24, 25). Purified spores ( $1 \times 10^7$ ) were suspended in 1 mL  
123 BHIS medium or germination salts [0.3 mM  $(\text{NH}_4)_2\text{SO}_4$ , 6.6 mM  $\text{KH}_2\text{PO}_4$ , 15 mM NaCl,  
124 59.5 mM  $\text{NaHCO}_3$ , and 35.2 mM  $\text{Na}_2\text{HPO}_4$ ] supplemented with 10 mM glycine. Cultures  
125 were further supplemented with 2 mM taurocholic acid (TA) and the test compounds at  
126 8X or 80X MIC. The initiation of spore germination was detected as the loss of optical  
127 density at 600 nm in a PerkinElmer Lambda 25 spectrophotometer.

128

129 **Measurements of killing rates.** For experiments involving phase-dark spores,  $1 \times 10^7$   
130 phase-bright spores of strain UK1 were first incubated for 10 minutes in BHIS medium  
131 supplemented with 2 mM TA. The suspension was then centrifuged for 1 min at 14,000  
132 x g. The resulting pellet of phase-dark spores was suspended in BHIS medium at a  
133 concentration of  $1 \times 10^5$  per ml with or without the test compounds at 8X MIC or 80X  
134 MIC in Teflon-coated tubes to prevent the adherence of spores to the tubes. Spores

135 were confirmed to be phase-dark by phase-contrast microscopy. For experiments using  
136 exponential phase vegetative cells, strains UK1 or JIR8094 ( $1 \times 10^5$  per ml) were  
137 suspended in BHIS medium supplemented with  $\text{CaCl}_2$  with or without addition of the  
138 test compounds at 8X MIC for strain JIR8094 or at 8X MIC and 80X MIC for strain UK1.  
139 For stationary phase cell killing assays, strain UK1 was grown in TY medium for ~12  
140 hours and then diluted in 30 ml TY supplemented with  $\text{CaCl}_2$  to give an  $\text{OD}_{600} \sim 0.1$ . After  
141 12 hrs incubation at  $37^\circ\text{C}$ , at which point the cells had left exponential growth phase,  
142 the culture was subdivided and the test compounds were each added to a separate  
143 culture tube at 8X or 80X MIC. For all assays, samples were withdrawn at indicated time  
144 points thereafter and viability was scored by plating serial dilutions on BHIS agar without  
145 calcium or antibiotic. Plates were incubated in the anaerobic chamber overnight. Each  
146 killing rate assay was performed at least twice to ensure reproducibility of the results.

147

148 **Measurements of toxin gene expression.** An overnight culture of strain UK1 was  
149 diluted in 30 ml TY- $\text{Ca}^{++}$  to give an  $\text{OD}_{600} \sim 0.1$ . After 12 hrs incubation at  $37^\circ\text{C}$ , the  
150 culture was split and each test compound was added to a separate subculture at 8X  
151 MIC. (Toxin gene expression was not tested in cells exposed to drugs at 80X MIC  
152 because such treatment with surotomycin at that concentration led to cell death and  
153 prevented the isolation of intact mRNA.) RNA was prepared from cells harvested at 0, 2,  
154 4, 8 and 24 hrs after addition of the drugs as previously described (26). RNA was  
155 quantified by absorbance using a NanoDrop ND-1000 spectrophotometer (Thermo  
156 Scientific). Primers for qPCR were designed using the online PrimerQuest tool from  
157 Integrated DNA Technologies

158 (<http://www.idtdna.com/Scitools/Applications/Primerquest>). Synthesis of cDNA was  
159 performed on 500 ng of RNA using random hexamer primers and the QuantiTect®  
160 Reverse Transcription Kit (QIAGEN) according to the manufacturer's recommendations.  
161 To control for chromosomal DNA contamination, mock cDNA synthesis reactions  
162 containing no reverse transcriptase were used as negative controls in subsequent  
163 amplifications. cDNA samples were diluted 4-fold and used as templates for qPCR of  
164 *rpoC* (primers oLB122 [CTAGCTGCTCCTATGTCTCACATC] and oLB123  
165 [CCAGTCTCTCCTGGATCAACTA]) and *tcdA* (primers oLB131  
166 [GTATGGATAGGTGGAGAAGTCA] and oLB132  
167 [CTCTTCCTCTAGTAGCTGTAATGC]) using the Roche SYBR Green I PCR mix and a  
168 Roche LightCycler 480 II thermocycler. Reactions were performed in a final volume of  
169 20 µl using 4 µl diluted cDNA and primers at 1 µM final concentration. Amplification  
170 included 45 cycles of the following steps: 10 s at 95 °C, 10 s at 52 °C, 15 s at 72 °C.  
171 Reactions were performed in triplicate using cDNA synthesized from each of a minimum  
172 of three biological replicates, and results are presented as the means and SEM of the  
173 data obtained. Results were calculated using the  $2^{-\Delta\Delta C_t}$  method, in which the  
174 amount of target mRNA is normalized to that of an internal control transcript (*rpoC*) (27).

175

176 **Measurements of toxin protein accumulation by ELISA.** Culture supernatants of the  
177 samples used for RNA extraction (above) were assayed for toxin A levels by ELISA (28).  
178 In brief, 96-well plates were coated with antibody to toxin A (Novus Biologicals)  
179 overnight at 4°C. After washing with PBS containing 0.1% Tween 20, the plates were  
180 incubated with blocking agent and, after discarding the agent, incubated with culture



181 fluid samples with or without dilution for 1 hr at room temperature. After washing with  
182 PBS-Tween, the plates were incubated with secondary antibodies fused to HRP (Gallus  
183 Immunotech, Inc.). After 1 hr at room temperature, the plates were washed again with  
184 PBS-Tween and mixed with peroxidase substrate for 40 min at room temperature. The  
185 reactions were stopped with  $\text{H}_2\text{SO}_4$  and the  $\text{OD}_{450}$  was determined using a plate reader.  
186 Standard curves were generated using purified recombinant toxin A (29).

187

## 188 RESULTS

189

190 **Determining the MICs of the test compounds.** The MICs for each of the test  
191 compounds were determined using exponential phase cells as the target. The MIC  
192 values varied according to the strain being tested and the growth medium used (**Table**  
193 **1**). For all further experimentation, the concentration of each antibiotic used was based  
194 on the MIC of that antibiotic in the strain to be tested growing in the medium being used.  
195 The growth medium chosen for each experiment was based on previous experimental  
196 conditions.

197

198 **MIC determination in mutant strains.** A panel of mutant strains derived from JIR8094  
199 and altered in susceptibility to nisin and other small, cationic antimicrobial peptides  
200 (CAMPs) was tested for their susceptibility to surotomycin, metronidazole and  
201 vancomycin. As shown in **Table 2**, all strains had identical MIC values for metronidazole,  
202 but the MIC values for surotomycin and vancomycin varied slightly. MC112 has a  
203 mutation in an autolysin that presumably alters the cell wall of the bacterium, resulting in

204 increased tolerance to nisin, though not a higher MIC value. (That is, MC112 survives  
205 exposure to higher concentrations of nisin than does the parent strain, but does not  
206 grow in the presence of nisin.) MC112 had similar growth to the wild-type parent in all  
207 antibiotics tested, including surotomycin, in which the strain grew at a 2-fold higher  
208 concentration.

209

210 MC119 carries a mutation in a regulatory protein that is known to control an ABC-  
211 transporter system (*cprABC*). As a result, the MC119 mutant has an extraordinarily high  
212 MIC value for growth in nisin. In a previous study, we determined that MC119 also is  
213 less susceptible to polymyxin B (16). The MC119 strain had a 2-fold higher MIC for  
214 both vancomycin and surotomycin. The strongly decreased susceptibility of MC119 to  
215 vancomycin and its modestly decreased susceptibility to surotomycin, and polymyxin B  
216 are consistent with the hypothesis that this mutant is altered in its response to  
217 antimicrobials that target the cell surface.

218

219 **Effects of antibiotic exposure on the initiation of *C. difficile* UK1 spore**  
220 **germination.** To test if surotomycin or vancomycin or metronidazole is able to affect *C.*  
221 *difficile* spore germination, purified spores of strain UK1 were incubated in BHIS-Ca<sup>++</sup>  
222 with or without the germinant taurocholate (TA) (10) and with or without each of the  
223 antibiotics. The spores germinated, as measured by loss of OD<sub>600</sub>, when suspended in  
224 BHIS-Ca<sup>++</sup> supplemented with 2 mM TA, but not in the absence of TA (**Figure 1A – D**).  
225 (The loss of OD corresponds to the conversion of spores from a birefringent [phase-  
226 bright] form to a non-birefringent [phase-dark] form.) The addition of surotomycin or

227 metronidazole or vancomycin at 8x or 80x MIC had no effect, either positive or negative,  
228 on TA-dependent spore germination (**Figure 1A – D**). When similar experiments were  
229 performed in a buffer containing TA and glycine, a co-germinant, the same lack of effect  
230 of the antibiotics was seen (data not shown).

231

232 **Effects of antibiotic exposure on germinated (phase-dark) spores.** To determine  
233 the time after spore germination at which outgrowing vegetative cells become sensitive  
234 to the antibiotics, we added antibiotics to germinated (phase-dark) spores in BHIS-Ca<sup>++</sup>  
235 and followed bacterial viability at intervals thereafter. When exposed to surotomycin, the  
236 outgrowing cells lost viability more rapidly than after exposure to the other drugs (**Table**  
237 **3**). Note that this experiment does not distinguish between the time needed for killing  
238 and the growth stage at which outgrowing vegetative cells become susceptible to the  
239 drugs.

240

241 **Effects of surotomycin, metronidazole and vancomycin on exponential-phase**  
242 **cells.** Strains UK1 and JIR8094 were grown at 37°C to early exponential phase in  
243 BHIS-Ca<sup>++</sup>. After determining the initial titer, the cultures were divided and inoculated  
244 with the test drugs at concentrations corresponding to 8X or 80X MIC. Samples of each  
245 culture were taken at multiple time points and surviving cell titers were determined by  
246 serial dilution and plating on BHIS-Ca<sup>++</sup> supplemented with taurocholic acid (TA). In  
247 exponential phase cells of strain UK1, surotomycin reduced viability by 90% more  
248 rapidly than did the other drugs (**Table 3**). In exponential phase cells of strain JIR8094,

249 surotomycin, metronidazole and nisin were all rapid killers, whereas vancomycin  
250 reduced viability more slowly (**Fig. 2**).

251

252 **Effects of surotomycin, metronidazole and vancomycin on stationary phase cells.**

253 After incubation of a culture of strain UK1 in TY + CaCl<sub>2</sub> at 37°C for 12 hrs, the culture  
254 was split and each drug was added to a separate culture tube at 8X or 80X MIC.  
255 Samples were harvested at various intervals for CFU counts and OD<sub>600</sub> readings. At 8X  
256 MIC, none of the three drugs reduced the turbidity of the culture (apparent cell mass) or  
257 viability substantially (**Figs. 3A and 4A**). At 80X MIC, however, surotomycin greatly  
258 diminished viability (but not cell mass significantly, indicating killing without cell lysis),  
259 whereas vancomycin and metronidazole had no detectable impact on cell mass or  
260 viability (**Figs. 3B and Fig. 4B**).

261

262 **Effects of surotomycin, metronidazole and vancomycin on toxin gene expression**

263 **and toxin release.** To test the effects of the various drugs on toxin production, cells at  
264 an early stage of stationary phase were exposed to the individual compounds at 8X MIC.  
265 At intervals thereafter, samples were harvested and RNA was extracted for quantitation  
266 by qRT-PCR. As shown in **Fig. 5**, expression of *tcdA*, the gene that encodes toxin A,  
267 increased with time to similar extents in untreated cells and cells treated with drugs.

268

269 To measure toxin A release into the culture medium, the supernatant fluids of the  
270 samples used above for RNA extraction were assayed for toxin A levels by ELISA (see  
271 Materials and Methods). Standard curves were generated using purified toxin A. As

272 shown in **Fig 6**, the amount of toxin A in the culture fluid was not significantly altered  
273 compared to the no-drug control by exposure to surotomycin or metronidazole at 8x  
274 MIC at early stationary phase. Addition of vancomycin, however, reduced toxin release  
275 at 24 hrs.

276

## 277 **DISCUSSION**

278

279 The results presented here show that surotomycin kills cells that are growing out of  
280 germinated spores, exponential-phase cells and stationary-phase cells (in the latter  
281 case at 80X MIC only) of strain UK1 more rapidly than does metronidazole or  
282 vancomycin. Surotomycin is also a more rapid killer of exponential-phase cells of strain  
283 JIR8094 than is vancomycin. None of the antibiotics tested was able to inhibit or  
284 activate *C. difficile* spore germination. This result seems to fit with the mechanisms of  
285 action of the tested antibiotics. The initiation of spore germination, a mostly enzymatic  
286 process, would be unaffected by antibiotics that need to be metabolized before  
287 becoming potent (metronidazole) or by antibiotics that inhibit new cell wall synthesis  
288 (vancomycin). Surotomycin leads to loss of the proton gradient, but does so by  
289 depolarizing the membrane without causing permeability of molecules as large as 671  
290 daltons (19). Such a mechanism might explain its lack of effect on the initiation of spore  
291 germination if the pores created are also not large enough to allow the escape of  $\text{Ca}^{++}$ -  
292 DPA from the spore core, a step necessary for the completion of germination (12, 13).  
293 Alternatively, surotomycin might be unable to penetrate the spore coat and cortex in  
294 order to gain access to the spore inner membrane.

295

296 Since outgrowing spores are metabolically active and synthesize new cell wall  
297 peptidoglycan, it is not surprising that such spores are susceptible to all of the  
298 antibiotics tested, although the heightened susceptibility of such cells to surotomycin  
299 was not predictable. The greater activity of surotomycin may be due to its ability to  
300 prevent the generation of a proton gradient needed for metabolic functions to resume.  
301 After the loss of dormancy, metabolism resumes in the core. Subsequently, a  
302 vegetative cell begins to grow from the germinated spore and this requires new cell wall  
303 synthesis. Metronidazole, which requires cellular metabolism for activity (30), inhibited  
304 growth in an intermediate time-frame with respect to surotomycin and vancomycin.  
305 Because new cell wall synthesis occurs late during outgrowth, it is not surprising that  
306 vancomycin required more time to inhibit growth than the other two antibiotics tested.

307

308 Although the ability of all three antibiotics to kill exponential phase cells was expected,  
309 their differential effects on stationary phase cells suggest that surotomycin may be more  
310 effective than vancomycin or metronidazole in reducing the severity and recurrence of  
311 infection. That is, at 80X MIC, a relatively low dose compared to those used clinically,  
312 surotomycin killed stationary phase cells rapidly. Since stationary phase cells are the  
313 ones that form spores and produce toxins, a drug that kills both exponential and  
314 stationary phase cells may be particularly effective in reducing the spore titer and toxin  
315 levels in the colon and in stool.

316

317 The appearance of *C. difficile* mutants that have decreased susceptibility to surotomycin  
318 is rare and the MIC shift is not more than 8-to-16 fold (18). Therefore, it is interesting  
319 that certain mutant strains that were isolated on the basis of their increased tolerance or  
320 resistance to nisin also showed slightly reduced susceptibility to surotomycin (**Table 2**).  
321 The basis for this decreased sensitivity is not known, but is likely to reflect changes in  
322 cell surface architecture. The decrease in susceptibility gained, however, would not be  
323 expected to influence the effectiveness of surotomycin *in vivo*.

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325

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332

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- 420

421 **FIGURE LEGENDS**

422 **Figure 1. Effect of antibiotic treatment on initiation of *C. difficile* spore**  
423 **germination.**

424 *C. difficile* UK1 spores were suspended in BHIS-Ca<sup>++</sup> medium (black) or medium  
425 supplemented with 8x MIC (A) or 80x MIC (B) surotomycin or 8x MIC (C) or 80x (D)  
426 vancomycin or metronidazole. Where indicated, the medium was supplemented with  
427 taurocholic acid (TA), an activator of germination. The initiation of spore germination  
428 was detected by measuring the loss of OD<sub>600</sub>. Experiments were performed in duplicate;  
429 the presented data represent one such experiment. Germination plots from the two  
430 experiments were superimposable.

431 **Figure 2. Effects of drugs at 8X MIC on survival of exponential phase cells of *C.***  
432 ***difficile* strain JIR8094.**

433 Exponential phase cultures (OD<sub>600</sub> = 0.45) in BHIS-Ca<sup>++</sup> were exposed to the three  
434 antibiotics and samples withdrawn at indicated times thereafter were assayed for  
435 viability (CFU/ml).

436 **Figure 3. Effects of drugs on stationary phase cells of strain UK1 as measured by**  
437 **cell mass.**

438 Panel A, each drug was added at 8X MIC to the cultures at early stationary phase;  
439 panel B, each drug was added at 80X MIC to the cultures at early stationary phase. At  
440 least 3 biological replicates were performed for each drug at each concentration.

441 **Figure 4. Effects of drugs at 8X and 80X MIC on viability of strain UK1 stationary**  
442 **phase cells.** Early stationary phase cells were exposed to antibiotics at 8X MIC (part A)

443 or 80X MIC (part B). At indicated time-points thereafter, samples were removed and  
444 plated for viable counts (CFU).

445 **Figure 5. qRT-PCR analysis of *tcdA* expression following treatment with**  
446 **surotomicin, metronidazole or vancomycin at 8X MIC.** Strain UK1 was grown in  
447 CaCl<sub>2</sub>-supplemented TY medium for ~12 hrs as described above, at which time the  
448 culture was split and drugs were added at 8X MIC. At two, four, eight and twenty-four  
449 hrs after addition of drugs, cell samples were harvested, RNA was extracted, and cDNA  
450 was synthesized as described under Materials and Methods. cDNA corresponding to  
451 *tcdA* mRNA was quantified by real time-PCR. Reactions were performed in triplicate  
452 using cDNA synthesized from each of a minimum of three biological replicates, and  
453 results are presented as the means and SEM of the data obtained. Results were  
454 calculated using the  $2^{-\Delta\Delta Ct}$  method, in which the amount of target mRNA was  
455 normalized to that of an internal control transcript (*rpoC*).

456 **Figure 6. ELISA assays of toxin A in cultures of strain UK1 treated with**  
457 **surotomicin, metronidazole or vancomycin at 8X MIC.** Culture fluids of cells  
458 exposed to antibiotics at early stationary phase were collected at the indicated times  
459 and assayed for toxin A by ELISA.

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466 **Table 1. MIC values for antimicrobial compounds<sup>a</sup>**

467

Strain (Growth Medium)	MIC (μg/ml)		
	Surotomycin	Metronidazole	Vancomycin
UK1 (BHIS + Ca <sup>++</sup> )	1.5	0.5	0.3125
JIR8094 (BHIS + Ca <sup>++</sup> )	1	0.5	2
UK1 (TY + Ca <sup>++</sup> )	0.125	0.125	1
ATCC 70057 (TY + Ca <sup>++</sup> )	0.125-0.25	0.06-0.125	1

468

469 <sup>a</sup> The various strains were grown in the indicated media and assayed for susceptibility  
470 to the indicated antibiotics provided in a series of two-fold dilutions. The MIC was  
471 defined as the lowest concentration of antibiotic that prevented measurable growth.

472

473 **Table 2. MIC values for antimicrobial compounds on mutant strains<sup>a</sup>**

474

Strain	MIC (μg/ml)			
	Nisin	Surotomycin	Metronidazole	Vancomycin
JIR8094	180	1	0.5	2
MC112	180	2	0.5	2
MC119	1440	2	0.5	4
MC120	90	1	0.5	2
MC141	90	1	0.5	2

483 <sup>a</sup> Each strain and each drug concentration was tested in cells growing in BHIS medium  
484 in duplicate for each assay. Assays were performed at least 3 times to ensure  
485 reproducibility of results.



486 **Table 3. Time required to reduce viability of *C. difficile* UK1 phase-dark spores and exponential-phase cells by**  
487 **90%.**

	Time to 90% Killing in Hours					
	8x MIC			80x MIC		
	Surotomycin	Metronidazole	Vancomycin	Surotomycin	Metronidazole	Vancomycin
UK1 phase-dark spores <sup>a</sup>	3.33	> 7	> 7	2	3	> 5
UK1 exponential-phase cells <sup>b</sup>	0.33	1.5	1.5	< 0.33	0.66	1.33

488 <sup>a</sup>Tubes containing BHIS medium supplemented with CaCl<sub>2</sub> and the indicated antibiotics (at 8X MIC or 80X MIC) were  
489 inoculated with phase-dark spores of strain UK1 and tested, after various times of incubation, for survival by plating on  
490 BHIS medium.

491 <sup>b</sup>Exponential-phase cells of strain UK1 growing in BHIS medium supplemented with CaCl<sub>2</sub> were exposed to two different  
492 concentrations (8X MIC and 80X MIC) of the indicated antibiotics. Survival was assayed by plating serial dilutions on  
493 BHIS-Ca<sup>++</sup> medium.

494

















